

Amendments to the Claims:

This listing of claims will replace all prior listings of claims in the application:

Listing of Claims:

1. (currently amended) A method of detecting a breast cancer cell in a biological sample from a patient, the method comprising
contacting the sample with a polynucleotide that selectively hybridizes to a nucleic acid sequence encoding a polypeptide having an amino acid sequence of SEQ ID NO:2, ~~SEQ ID NO:4, or SEQ ID NO:6~~; and
detecting an increase in the level of the nucleic acid sequence, relative to normal, thereby detecting the presence of a breast cancer in the patient.
2. (original) The method of claim 1, wherein the detecting step comprises detecting an mRNA that encodes the polypeptide.
3. (original) The method of claim 2, wherein the mRNA is detected using an amplification reaction.
4. (original) The method of claim 1, wherein the detecting step comprises detecting an increase in copy number of the nucleic acid that encodes the polypeptide.
5. (original) The method of claim 1, wherein the patient is undergoing a therapeutic regimen to treat breast cancer.
6. (original) The method of claim 1, wherein the patient is suspected of having breast cancer.
7. (original) A method of detecting a breast cancer cell in a biological sample from a patient, the method comprising

detecting an increase in the level of a polypeptide having an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4, or SEQ ID NO:6, relative to normal, thereby detecting the presence of a breast cancer in the patient.

8. (original) The method of claim 7, wherein the step of detecting an increase in the level of the polypeptide comprises performing an immunoassay.

9. (original) A method of monitoring the efficacy of a therapeutic treatment of cancer, the method comprising the steps of:

(i) providing a biological sample from a patient undergoing the therapeutic treatment; and

(ii) detecting the level of: a polypeptide having an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4, or SEQ ID NO:6, or of a nucleic acid that encodes the polypeptide, in the biological sample compared to a level in a biological sample from the patient prior to, or earlier in, the therapeutic treatment, thereby monitoring the efficacy of the therapy.

10. (original) A method for identifying a compound that modulates a breast cancer-associated polypeptide, the method comprising the steps of:

(i) contacting the compound with a polypeptide of SEQ ID NO:2, SEQ ID NO:4, or SEQ ID NO:6; and

(ii) determining the functional effect of the compound upon the polypeptide.

11. (original) A method of inhibiting proliferation of a breast cancer cell that overexpresses a polypeptide having an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4, or SEQ ID NO:6, the method comprising the step of contacting the cancer cell with a therapeutically effective amount of an inhibitor of the polypeptide.

12. (original) The method of claim 11, wherein the gene that encodes the polypeptide is increased in copy number in the breast cancer cell.

13. (original) The method of claim 11, wherein the inhibitor is an antibody.
14. (original) The method of claim 11, wherein the inhibitor is a small molecule.
15. (new) The method of claim 1, wherein the biological sample is breast tissue.